

GenCore version 6.3  
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:19:14 ; Search time 581 Seconds  
(without alignments)  
14891.617 Million cell

updates/sec

Title: US-09-300-482-619  
Perfect score: 300  
Sequence: 1  
gatttattgacaacaccgat.....ttcagaaatgtctgctgtgg 300

Scoring table: IDENTITY\_NUC  
Gapop 10.0 , Gapext 1.0

Searched: 41078765 seqs, 14427166270 residues

Total number of hits satisfying chosen parameters: 82157530

Minimum DB seq length: 0  
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%  
Maximum Match 100%  
Listing first 45 summaries

Database : Published\_Applications\_NA\_Main:\*  
1:  
/ABSS/Data/CRF/ptodata/1/pubpna/US07\_PUBCOMB.seq:\*  
2:  
/ABSS/Data/CRF/ptodata/1/pubpna/US08\_PUBCOMB.seq:\*  
3:  
/ABSS/Data/CRF/ptodata/1/pubpna/US09A\_PUBCOMB.seq:\*  
4:  
/ABSS/Data/CRF/ptodata/1/pubpna/US09B\_PUBCOMB.seq:\*  
5:  
/ABSS/Data/CRF/ptodata/1/pubpna/US09C\_PUBCOMB.seq:\*  
6:  
/ABSS/Data/CRF/ptodata/1/pubpna/US09D\_PUBCOMB.seq:\*  
7:  
/ABSS/Data/CRF/ptodata/1/pubpna/US10A\_PUBCOMB.seq:\*  
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